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11/14

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## RAW SEQUENCE LISTING

DATE: 11/21/2002

PATENT APPLICATION: US/09/780,438B

TIME: 10:08:29

Input Set : A:\EP.txt

Output Set: N:\CRF4\11212002\I780438B.raw

3 <110> APPLICANT: Qi, Xiaoyang

5 <120> TITLE OF INVENTION: Fusogenic Properties of Saposin C and Related Proteins and Polypeptides

6 for Application to Transmembrane Drug Delivery Systems

8 <130> FILE REFERENCE: 60/181,754

10 <140> CURRENT APPLICATION NUMBER: US 09/780,438B

11 <141> CURRENT FILING DATE: 2001-02-09

13 <150> PRIOR APPLICATION NUMBER: 60/181,754

14 <151> PRIOR FILING DATE: 2000-02-11

16 <160> NUMBER OF SEQ ID NOS: 4

18 <170> SOFTWARE: PatentIn version 3.1

20 <210> SEQ ID NO: 1

21 <211> LENGTH: 38

22 <212> TYPE: PRT

23 <213> ORGANISM: Homo sapiens

25 <220> FEATURE:

26 <221> NAME/KEY: MISC\_FEATURE

27 <222> LOCATION: (1)..(1)

28 <223> OTHER INFORMATION: Where the amino acid located at 1 is a hydrophobic amino acid,

29 including Val, Leu, Ile, Met, Pro, Phe, and Ala

32 <220> FEATURE:

33 <221> NAME/KEY: MISC\_FEATURE

34 <222> LOCATION: (2)..(2)

35 <223> OTHER INFORMATION: Where the amino acid located at 2 is an uncharged polar amino acid,

36 including Thr, Ser, Tyr, Gly, Gln, and Asn

39 <220> FEATURE:

40 <221> NAME/KEY: MISC\_FEATURE

41 <222> LOCATION: (5)..(5)

42 <223> OTHER INFORMATION: Where the amino acid located at 5 is a hydrophobic amino acid,

43 including Val, Leu, Ile, Met, Pro, Phe, and Ala

46 <220> FEATURE:

47 <221> NAME/KEY: MISC\_FEATURE

48 <222> LOCATION: (8)..(10)

49 <223> OTHER INFORMATION: Where the amino acids located at 8-10 are hydrophobic amino acids,

50 including Val, Leu, Ile, Met, Pro, Phe, and Ala

53 <220> FEATURE:

54 <221> NAME/KEY: MISC\_FEATURE

55 <222> LOCATION: (13)..(13)

56 <223> OTHER INFORMATION: Where the amino acid located at 13 is a hydrophobic amino acid,

ENTERED

57 including Val, Leu, Ile, Met, Pro, Phe, and Ala  
59 <220> FEATURE:  
60 <221> NAME/KEY: MISC\_FEATURE  
61 <222> LOCATION: (14)..(14)  
62 <223> OTHER INFORMATION: Where the amino acid located at 14 is an uncharged polar  
amino acid,

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63 including Thr, Ser, Tyr, Gly, Gln, and Asn  
 66 <220> FEATURE:  
 67 <221> NAME/KEY: MISC\_FEATURE  
 68 <222> LOCATION: (16)..(17)  
 69 <223> OTHER INFORMATION: Where the amino acids located at 16 and 17 are hydrophobic  
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 70 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 73 <220> FEATURE:  
 74 <221> NAME/KEY: MISC\_FEATURE  
 75 <222> LOCATION: (22)..(22)  
 76 <223> OTHER INFORMATION: Where the amino acid located at 22 is an uncharged polar  
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 77 acid, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 80 <220> FEATURE:  
 81 <221> NAME/KEY: MISC\_FEATURE  
 82 <222> LOCATION: (26)..(27)  
 83 <223> OTHER INFORMATION: Where the amino acids located at 26 and 27 are hydrophobic  
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 84 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 87 <220> FEATURE:  
 88 <221> NAME/KEY: MISC\_FEATURE  
 89 <222> LOCATION: (29)..(30)  
 90 <223> OTHER INFORMATION: Where the amino acids located at 29 and 30 are hydrophobic  
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 91 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 94 <220> FEATURE:  
 95 <221> NAME/KEY: MISC\_FEATURE  
 96 <222> LOCATION: (33)..(33)  
 97 <223> OTHER INFORMATION: Where the amino acid located at 33 is a hydrophobic amino  
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 98 including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 101 <220> FEATURE:  
 102 <221> NAME/KEY: MISC\_FEATURE  
 103 <222> LOCATION: (35)..(35)  
 104 <223> OTHER INFORMATION: Where the amino acid located at 35 is an uncharged polar  
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 105 acid, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 108 <220> FEATURE:  
 109 <221> NAME/KEY: MISC\_FEATURE  
 110 <222> LOCATION: (37)..(38)  
 111 <223> OTHER INFORMATION: Where the amino acids located at 37 and 38 are hydrophobic  
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 112 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 115 <400> SEQUENCE: 1  
 W--> 117 Xaa Xaa Cys Glu Xaa Cys Glu Xaa Xaa Xaa Lys Glu Xaa Xaa Lys Xaa  
 118 1 5 10 15  
 W--> 121 Xaa Asp Asn Asn Lys Xaa Glu Lys Glu Xaa Xaa Asp Xaa Xaa Asp Lys  
 122 20 25 30  
 W--> 125 Xaa Cys Xaa Lys Xaa Xaa  
 126 35  
 129 <210> SEQ ID NO: 2  
 130 <211> LENGTH: 39  
 131 <212> TYPE: PRT

132 <213> ORGANISM: Homo sapiens  
134 <220> FEATURE:  
135 <221> NAME/KEY: MISC\_FEATURE

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136 <222> LOCATION: (1)..(2)

137 <223> OTHER INFORMATION: Where the amino acids located at 1 and 2 are hydrophobic amino acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala

141 <220> FEATURE:

142 <221> NAME/KEY: MISC\_FEATURE

143 <222> LOCATION: (3)..(3)

144 <223> OTHER INFORMATION: Where the amino acid located at 3 is an uncharged polar amino acid, including Thr, Ser, Tyr, Gly, Gln, and Asn

148 <220> FEATURE:

149 <221> NAME/KEY: MISC\_FEATURE

150 <222> LOCATION: (6)..(6)

151 <223> OTHER INFORMATION: Where the amino acid located at 6 is a hydrophobic amino acid, including Val, Leu, Ile, Met, Pro, Phe, and Ala

155 <220> FEATURE:

156 <221> NAME/KEY: MISC\_FEATURE

157 <222> LOCATION: (9)..(11)

158 <223> OTHER INFORMATION: Where the amino acids located at 9-11 are hydrophobic amino acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala

162 <220> FEATURE:

163 <221> NAME/KEY: MISC\_FEATURE

164 <222> LOCATION: (14)..(14)

165 <223> OTHER INFORMATION: Where the amino acid located at 14 is a hydrophobic amino acid, including Val, Leu, Ile, Met, Pro, Phe, and Ala

169 <220> FEATURE:

170 <221> NAME/KEY: MISC\_FEATURE

171 <222> LOCATION: (15)..(15)

172 <223> OTHER INFORMATION: Where the amino acid located at 15 is an uncharged polar amino acid, including Thr, Ser, Tyr, Gly, Gln, and Asn

176 <220> FEATURE:

177 <221> NAME/KEY: MISC\_FEATURE

178 <222> LOCATION: (17)..(18)

179 <223> OTHER INFORMATION: Where the amino acids located at 17 and 18 are hydrophobic amino acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala

183 <220> FEATURE:

184 <221> NAME/KEY: MISC\_FEATURE

185 <222> LOCATION: (23)..(23)

186 <223> OTHER INFORMATION: Where the amino acid located 23 is an uncharged polar amino acid, including Thr, Ser, Tyr, Gly, Gln, and Asn

190 <220> FEATURE:

191 <221> NAME/KEY: MISC\_FEATURE

192 <222> LOCATION: (27)..(28)

193 <223> OTHER INFORMATION: Where the amino acids located at 27 and 28 are hydrophobic amino acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala

197 <220> FEATURE:

198 <221> NAME/KEY: MISC\_FEATURE  
199 <222> LOCATION: (30)..(31)  
200 <223> OTHER INFORMATION: Where the amino acids located at 30 and 31 are hydrophobic  
amino  
201 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
204 <220> FEATURE:

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Input Set : A:\EP.txt

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205 <221> NAME/KEY: MISC\_FEATURE  
 206 <222> LOCATION: (34)..(34)  
 207 <223> OTHER INFORMATION: Where the amino acid located at 34 is a hydrophobic amino  
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 208 including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 211 <220> FEATURE:  
 212 <221> NAME/KEY: MISC\_FEATURE  
 213 <222> LOCATION: (36)..(36)  
 214 <223> OTHER INFORMATION: Where the amino acid located at 36 is an uncharged polar  
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 215 acid, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 218 <220> FEATURE:  
 219 <221> NAME/KEY: MISC\_FEATURE  
 220 <222> LOCATION: (38)..(39)  
 221 <223> OTHER INFORMATION: Where the amino acids located at 38 and 39 are hydrophobic  
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 222 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 225 <400> SEQUENCE: 2  
 W--> 227 Xaa Xaa Xaa Cys Glu Xaa Cys Glu Xaa Xaa Xaa Lys Glu Xaa Xaa Lys  
 228 1 5 10 15  
 W--> 231 Xaa Xaa Asp Asn Asn Lys Xaa Glu Lys Glu Xaa Xaa Asp Xaa Xaa Asp  
 232 20 25 30  
 W--> 235 Lys Xaa Cys Xaa Lys Xaa Xaa  
 236 35  
 239 <210> SEQ ID NO: 3  
 240 <211> LENGTH: 38  
 241 <212> TYPE: PRT  
 242 <213> ORGANISM: Homo sapiens  
 244 <220> FEATURE:  
 245 <221> NAME/KEY: MISC\_FEATURE  
 246 <222> LOCATION: (1)..(1)  
 247 <223> OTHER INFORMATION: Where the amino acid located at 1 is a hydrophobic amino  
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 248 including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 251 <220> FEATURE:  
 252 <221> NAME/KEY: MISC\_FEATURE  
 253 <222> LOCATION: (2)..(2)  
 254 <223> OTHER INFORMATION: Where the amino acid located at 2 is an uncharged polar  
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 255 including Thr, Ser, Tyr, Gly, Gln, and Asn  
 258 <220> FEATURE:  
 259 <221> NAME/KEY: MISC\_FEATURE  
 260 <222> LOCATION: (5)..(5)  
 261 <223> OTHER INFORMATION: Where the amino acid located at 5 is a hydrophobic amino  
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 262 including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 265 <220> FEATURE:  
 266 <221> NAME/KEY: MISC\_FEATURE  
 267 <222> LOCATION: (8)..(10)  
 268 <223> OTHER INFORMATION: Where the amino acids located at 8-10 are hydrophobic amino  
 acids,  
 269 including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 272 <220> FEATURE:

273 <221> NAME/KEY: MISC\_FEATURE

274 <222> LOCATION: (13)..(13) ✓

275 <223> OTHER INFORMATION: Where the amino acid located at 13 is a hydrophobic amino acid,

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276 including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 279 <220> FEATURE:  
 280 <221> NAME/KEY: MISC\_FEATURE  
 281 <222> LOCATION: (14)..(14)  
 282 <223> OTHER INFORMATION: Where the amino acid located at 14 is an uncharged polar  
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 283 acid, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 286 <220> FEATURE:  
 287 <221> NAME/KEY: MISC\_FEATURE  
 288 <222> LOCATION: (16)..(17)  
 289 <223> OTHER INFORMATION: Where the amino acids located at 16 and 17 are hydrophobic  
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 290 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 293 <220> FEATURE:  
 294 <221> NAME/KEY: MISC\_FEATURE  
 295 <222> LOCATION: (22)..(22)  
 296 <223> OTHER INFORMATION: Where the amino acid located at 22 is an uncharged polar  
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 297 acid, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 300 <220> FEATURE:  
 301 <221> NAME/KEY: MISC\_FEATURE  
 302 <222> LOCATION: (26)..(27)  
 303 <223> OTHER INFORMATION: Where the amino acids located at 26 and 27 are hydrophobic  
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 304 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 307 <220> FEATURE:  
 308 <221> NAME/KEY: MISC\_FEATURE  
 309 <222> LOCATION: (29)..(30)  
 310 <223> OTHER INFORMATION: Where the amino acids located at 29 and 30 are hydrophobic  
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 311 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 314 <220> FEATURE:  
 315 <221> NAME/KEY: MISC\_FEATURE  
 316 <222> LOCATION: (33)..(33)  
 317 <223> OTHER INFORMATION: Where the amino acid located at 33 is a hydrophobic amino  
 acid,  
 318 including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 321 <220> FEATURE:  
 322 <221> NAME/KEY: MISC\_FEATURE  
 323 <222> LOCATION: (35)..(35)  
 324 <223> OTHER INFORMATION: Where the amino acid located at 35 is an uncharged polar  
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 325 acid, including Thr, Ser, Tyr, Gly, Gln, and Asn  
 328 <220> FEATURE:  
 329 <221> NAME/KEY: MISC\_FEATURE  
 330 <222> LOCATION: (37)..(38)  
 331 <223> OTHER INFORMATION: Where the amino acids located at 37 and 38 are hydrophobic  
 amino  
 332 acids, including Val, Leu, Ile, Met, Pro, Phe, and Ala  
 335 <400> SEQUENCE: 3  
 W--> 337 Xaa Xaa Cys Glu Xaa Cys Glu Xaa Xaa Xaa Lys Glu Xaa Xaa Lys Xaa  
 338 1 5 10 15  
 W--> 341 Xaa Asp Asn Asn Lys Xaa Glu Lys Glu Xaa Xaa Asp Xaa Xaa Asp Lys

342	20	25	30
W--> 345	Xaa Cys Xaa Lys Xaa Xaa		
346	35		
349	<210> SEQ ID NO: 4		

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/780,438B

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Input Set : A:\EP.txt

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1, 2, 5, 8, 9, 10, 13, 14, 16, 17, 22, 26, 27, 29, 30, 33, 35, 37, 38

Seq#:2; Xaa Pos. 1, 2, 3, 6, 9, 10, 11, 14, 15, 17, 18, 23, 27, 28, 30, 31, 34, 36, 38, 39

Seq#:3; Xaa Pos. 1, 2, 5, 8, 9, 10, 13, 14, 16, 17, 22, 26, 27, 29, 30, 33, 35, 37, 38

Seq#:4; Xaa Pos. 1, 2, 5, 8, 9, 10, 13, 14, 16, 17, 22, 26, 27, 29, 30, 33, 35, 37, 38

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/780,438B

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Input Set : A:\EP.txt

Output Set: N:\CRF4\11212002\I780438B.raw

L:117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0  
L:121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:16  
L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:32  
L:227 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0  
L:231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:16  
L:235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:32  
L:337 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0  
L:341 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:16  
L:345 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:32  
L:449 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0  
L:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:16  
L:457 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:32